

Kaufman

CRF Errors Corrected by the STIC System Branch

1646 #9

Serial Number: 09/021,660

CRF Processing Date: 7/20/99
Edited by: [Signature]
Verified by: [Signature] (STIC staff)

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: **ENTERED**
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;
☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

PAGE: 1

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/021,660DATE: 07/20/1999
TIME: 16:25:59

Input Set: I021660.RAW

This Raw Listing contains the General
Information Section and those Sequences
containing ERRORS.

1 <110> Harvard University
2 Baron, Margaret H. et al.
3 <120> Methods for Modulating Hematopoiesis and
4 Vascular Growth
5 <130> 1874/110
6 <140> US/09/021,660
7 <141> 1998-02-10
8 <150> 60/037,513
9 <151> 1997-02-10
10 <150> 60/049,763
11 <151> 1997-06-16
12 <160> 26
13 <170> FastSEQ for Windows Version 3.0

Does Not Comply
Corrected Diskette Needed

ERRORED SEQUENCES FOLLOW

14 <210> 26
15 <211> 20
16 <212> DNA
17 <213> PCR Primer
18 <400> 26
19 aagaaggata agaggacagg
E--> 20 63

20

PAGE: 2

VERIFICATION SUMMARY
PATENT APPLICATION US/09/021,660

DATE: 07/20/1999
TIME: 16:25:59

Input Set: I021660.RAW

Line ? Error/Warning

Original Text

20 E Number of Bases conflict w/ Running Total 63